

SEQUENCE LISTING

<110> COHEN, STEPHEN
BOUWMEESTER, ANTONIUS
ROYET, JULIEN

<120> REGULATOR OF NOTCH SIGNALING ACTIVITY

<130> 55880(71745)

<140> 09/830,980

<141> 2001-05-02

<150> PCT/IB99/01891

<151> 1999-11-03

<150> GB 9824045.0

<151> 1998-11-03

<160> 16

<170> PatentIn Ver. 2.1

<210> 1

<211> 480

<212> PRT

<213> Drosophila sp.

<400> 1

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Pro Ala Gly Ile Thr Thr Gln Gln Leu Gly Leu Ile Cys Asn Ala Leu
      35              40              45

Leu Lys Asn Glu Glu Ala Thr Pro Tyr Leu Phe Phe Val Gly Glu Asp
      50              55              60

Glu Ile Lys Lys Ser Leu Glu Asp Thr Leu Asp Leu Ala Ser Val Asp
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Thr Glu Asn Val Ile Asp Ile Val Tyr Gln Pro Gln Ala Val Phe Lys
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Val Arg Pro Val Thr Arg Cys Thr Ser Ser Met Pro Gly His Ala Glu
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Ala Val Val Ser Leu Asn Phe Ser Pro Asp Gly Ala His Leu Ala Ser
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Gly Ser Gly Asp Thr Thr Val Arg Leu Trp Asp Leu Asn Thr Glu Thr
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2004-10-08 09:00:00

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Ala 385	Ser	Phe	Asp	Lys	Ser 390	Val	Arg	Leu	Trp	Arg 395	Ala	Ser	Asp	Gly	Gln 400
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Trp	Ser	Ala	Asp 420	Ser	Arg	Leu	Ile	Val 425	Ser	Gly	Ser	Lys	Asp 430	Ser	Thr
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<213> *Drosophila* sp.

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<213> *Saccharomyces cerevisiae*

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Ser Ile Lys Phe Gln Ala Leu Asp Thr Gly Asp Asn Val Gly Gly Ala
 35 40 45

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 Val Ser Ala Ser Phe Asp Asn Ser Ile Lys Leu Trp Asp Gly Arg Asp
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 Gly Lys Phe Ile Ser Thr Phe Arg Gly His Ile Ala Ser Val Tyr Gln
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 Val Ala Trp Ser Ser Asp Cys Arg Leu Leu Val Ser Cys Ser Lys Asp
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 Thr Thr Leu Lys Val Trp Asp Val Arg Thr Arg Lys Leu Ser Val Asp
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 <213> Codonanthe elegans

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 Asn Gln Leu Leu Gly Ser Arg Phe Cys Leu Asn Asn Glu Phe Ser Val
 35 40 45
 Ser Gly Ala Glu Ile Val Asp Ser Ile Arg Lys Ser Leu Glu Glu Ile
 50 55 60

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 Pro Val Ile Ser Ala Gln Phe Ser Pro Asp Gly Arg Gly Leu Ala Ser
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 Gly Ser Gly Asp Gln Thr Met Arg Ile Trp Asp Ile Glu Leu Glu Leu
 115 120 125
 Pro Leu His Thr Cys Lys Ser His Lys Ser Trp Val Leu Cys Ile Ala
 130 135 140
 Trp Ser Pro Asp Ala Thr Lys Ile Ala Ser Ala Cys Lys Asn Gly Glu
 145 150 155 160
 Ile Cys Ile Trp Asn Ala Lys Thr Gly Glu Gln Ile Gly Lys Thr Leu
 165 170 175
 Lys Arg His Lys Gln Trp Ile Xaa Xaa Leu Ala Trp Gln Pro Thr Val
 180 185 190
 Lys Met Trp Arg Ala Asp Asp Gly Val Met Cys Arg Asn Met Thr Gly
 195 200 205
 His Ala His Trp Ile Asn Thr Leu Ala Leu Asn Thr Asp Tyr Ala Leu
 210 215 220
 Arg Thr Ser Cys Phe Glu Pro Ser Lys Ile Asn Arg Met Thr Gly His
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 Met Gln Leu Val Asn Gln Val Val Phe Ser Pro Asp Thr Arg Tyr Ala
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 Ser Ala Ser Phe Asp Lys Ser Val Lys Leu Trp Cys Gly Arg Thr Gly
 260 265 270
 Lys Tyr Leu Ala Ser Phe Arg Gly His Val Gly Pro Val Tyr Gln Val
 275 280 285
 Ala Trp Ser Ala Asp Ser Arg Leu Leu Val Ser Gly Ser Ala Asp Ser
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 Thr Leu Lys Val Phe Glu Leu Lys Thr Lys Ser Leu Tyr Tyr Asp Leu
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<211> 103

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35 40 45
Glu Glu Pro Leu Pro Leu Ala Phe Tyr Val His Asp Ala Glu Ile Val
50 55 60
Ser Ser Leu Gly Lys Thr Leu Glu Ser Gln Ser Val Glu Thr Glu Lys
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35 40 45
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Asp	Lys	Leu	Gln	Leu	Val	Cys	Asn	Ala	Leu	Leu	Gln	Glu	Glu	Asp	Pro	
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Val	Pro	Leu	Ala	Phe	Phe	Val	Gln	Asp	Leu	Glu	Ile	Val	Thr	Ser	Leu	
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Asp	Lys	Thr	Leu	Glu	Lys	Gln	Ser	Val	Glu	Thr	Glu	Lys	Val	Ile	Asp	
65					70					75					80	
Ile	Ile	Tyr	Gln	Pro	Gln	Ala	Val	Phe	Lys	Val	Arg	Ala	Val	Thr	Arg	
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Cys	Thr	Ser	Ser	Leu	Glu	Gly	His	Thr	Glu	Ala	Val	Ile	Ser	Val	Ala	
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Phe	Ser	Pro	Thr	Gly	Lys	Tyr	Leu	Ala	Ser	Gly	Ser	Gly	Asp	Thr	Thr	
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Lys	Leu	Ala	Ser	Gly	Cys	Lys	Asn	Ser	Gln	Ile	Phe	Ile	Trp	Asp	Pro	
				165					170					175		
Ser	Thr	Gly	Lys	Gln	Ile	Gly	Lys	Pro	Leu	Thr	Gly	His	Ser	Lys	Trp	
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	210					215					220					
Val	Met	Gly	Gln	Cys	Gln	Lys	Ile	Leu	Thr	Ser	His	Thr	Gln	Ser	Val	
225					230					235					240	
Thr	Ala	Val	Lys	Trp	Gly	Gly	Asp	Gly	Leu	Leu	Tyr	Ser	Ser	Ser	Gln	
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